## **Amendment To The Claims**

The following listing of claims will replace all prior versions, and listings, of claims in the application.

- 1. (Currently amended) A method for screening a candidate compound for effectiveness in modifying the binding properties interaction between ef a p38 protein and a parkan protein and comprising the steps of:
  - exposing a sample comprising said p38 protein and parkan protein to said candidate compound; and
- b) measuring a the <u>interaction between binding property</u> said p38 protein <u>to said</u> <u>parkan protein and comparing it to a control sample comprising said p38 protein and parkan protein without said candidate compound.</u>
- 2. (Canceled)
- 3. (Canceled)
- 4. (Original) The method of Claim 1 wherein said sample further comprises an enzyme substrate.
- 5. (Original) The method of Claim 1, wherein said method is performed in vitro.
- 6. (Canceled)
- 7. (Currently amended) The method of Claim 5, wherein said p38 protein is expressed in yeast.
- 8. (Currently amended) The method of Claim 1, wherein said p38 protein is selected from the group consisting of:
  - a) a polypeptide encoded by the polynucleotide of SEQ ID NO:1,
  - a polypeptide comprising the amino acid sequence of SEQ ID NO:2,
  - a polypeptide comprising the amino acid sequence of polynucleotide of SEQ ID NO:3,

- d) c) a polypeptide encoded by the polynucleotide of SEQ ID NO:4,
- e) d) a polypeptide having at least 65% identity to a polypeptide encoded by the polynucleotide of SEQ ID NO:1,
- f) e) a polypeptide having at least 65% identity to a polypeptide comprising the amino acid sequence of SEQ ID NO:2,
- g) a polypeptide having at least 65% identity to a polypeptide comprising the amino acid sequence of SEQ ID NO:3, and
- h) <u>f)</u> a polypeptide having at least 65% identity to a polypeptide encoded by the polynucleotide of SEQ ID NO:4.
- 9. (Currently amended) The method of Claim 2 1, wherein said parkin is selected from the group consisting of:
  - a) a polypeptide comprising the amino acid sequence of SEQ ID NO:5,
  - b) a polypeptide comprising the amino acid sequence of SEQ ID NO:6,
  - e) a) a polypeptide encoded by the polynucleotide of SEQ ID NO:7,
  - d) b) a polypeptide comprising the amino acid sequence of SEQ ID NO:8,
  - e) a polypeptide comprising the amine acid sequence of SEQ ID NO:9,
  - f) c) a polypeptide encoded by the polynucleotide of SEQ ID NO:10,
  - g) a polypeptide having at least 65% identity to a polypeptide comprising the amino acid sequence of SEQ ID NO:5,
  - h) a polypeptide having at least 65% identity to a polypeptide comprising the amino acid sequence of SEQ ID NO:6,
  - i) d) a polypeptide having at least 65% identity to a polypeptide encoded by the polynucleotide of SEQ ID NO:7,
  - <u>e)</u> a polypeptide having at least 65% identity to a polypeptide comprising the amino acid sequence of SEQ ID NO:8,
  - k) a polypeptide having at least 65% identity to a polypeptide comprising the amino acid sequence of SEQ ID NO:9, and
  - 4) <u>f)</u> a polypeptide having at least 65% identity to a polypeptide encoded by the polynucleotide of SEQ ID NO:10.
- 10. (Withdrawn) A compound identified accord to according to the method of Claim 1.
- 11. (Withdrawn) The compound of Claim 10, wherein said compound is selected from the group consisting of: a polypeptide, a polynucleotide, a lipid, a saccharide, and an antibody.

- 12. (Withdrawn) A pharmaceutical composition comprising an effective amount of the compound of Claim 10 and a pharmaceutically acceptable excipient.
- 13. (Withdrawn) A method of treating a neurodegenerative disease comprising the step of administering the pharmaceutical compound of Claim 12.
- 14. (Withdrawn) The method of Claim 13, wherein said neurodegenerative disease is Parkinson's disease.
- 15. (Withdrawn) An isolated polypeptide comprising an amino acid sequence selected from the group consisting of:
  - a) a polypeptide comprising an amino acid sequence of SEQ ID NO:2,
  - a polypeptide comprising an amino acid sequence comprising at least 5 consecutive amino acid residues of SEQ ID NO:2,
  - a polypeptide comprising an amino acid sequence comprising at least 9
    consecutive amino acid residues of SEQ ID NO:2,
  - d) a polypeptide comprising an amino acid sequence comprising at least 15 consecutive amino acid residues of SEQ ID NO:2,
  - e) a polypeptide comprising an amino acid sequence that is a derivative of SEQ ID NO:2, and
  - f) a polypeptide comprising an amino acid sequence that is a fragment of SEQ
    ID NO:2.
- 16. (Withdrawn) An isolated polynucleotide encoding a polypeptide of Claim 15.
- 17. (Withdrawn) The isolated polynucleotide of Claim 16 comprising the sequence of SEQ ID NO:1.
- 18. (Withdrawn) A vector comprising the isolated polynucleotide of Claim 16.
- 19. (Withdrawn) The vector of Claim 18, wherein said vector is a defective recombinant virus.
- 20. (Withdrawn) An isolated antibody which specifically binds to a polypeptide of claim 15.

- 21. (Withdrawn) The antibody of claim 20, wherein the antibody is selected from the group consisting of: a chimeric antibody, a single chain antibody, a Fab fragment, a  $\frac{F(ab').sub.2}{F(ab')_2}$  fragment, and a humanized antibody.
- 22. (Withdrawn) A method for producing a polypeptide of claim 15, the method comprising:
  - a) culturing a cell under conditions suitable for expression of the polypeptide, wherein said cell is transformed with a recombinant polynucleotide, and said recombinant polynucleotide comprises a promoter sequence operably linked to a polynucleotide encoding the polypeptide of claim 15, and
  - b) recovering the polypeptide so expressed.
- 23. (Withdrawn) A cell transformed with a recombinant polynucleotide of Claim 16.

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